

# Temporal dynamics of antibiotics in wastewater treatment plant influent

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## Abstract

A yearlong field experimental campaign was conducted to reveal time scales over which antibiotic fluxes vary in the influent of a wastewater treatment plant (WTP). In particular, sampling was carried out to ascertain the amplitudes of monthly, daily and hourly fluctuations of several antibiotics. A total of 180 samples was collected at the entrance of a WTP in Lausanne, Switzerland. Sample concentrations were multiplied by flow rate to obtain monthly, daily and hourly mass fluxes of six antibiotics (trimethoprim, norfloxacin, ciprofloxacin, ofloxacin, clindamycin and metronidazole). Seasonality in mass fluxes was observed for all substances, with maximum values in winter being up to an order of magnitude higher than in summer. The hourly measurements of the mass flux of antibiotics were found to have a period of 12 h. This was due to peaks in toilet use in the morning and early evening. In particular, the morning peak in flushing coincided with high concentrations (and hence high mass fluxes) due to overnight accumulation of substances

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in urine. However, little variation was observed in the average daily flux. Consequently, fluctuations in mass fluxes of antibiotics were mainly evident at the monthly and hourly time scales, with little variation on the day-week time scale. These results can aid in optimizing removal strategies and future sampling campaigns focused on antibiotics in wastewater.

*Keywords:* Pharmaceuticals, Water quality, Mass loading, Sampling, Time scale, Fluctuation

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## 1 **1. Introduction**

2     Although the quality of natural waters has been investigated extensively,  
3 continuing efforts are directed to understand better the complex dynamics  
4 of wastewater discharges to the environment. The most studied and best  
5 treated wastewater pollutants are organic materials and nutrients. On the  
6 other hand, the origin and fate of emerging pollutants, also referred as mi-  
7 cropollutants, are still insufficiently understood (Ternes and Joss, 2006; Ver-  
8 licchi et al., 2012). There is a plethora of micropollutants, each of which  
9 potentially displays different behavior in the environment (Kleywegt et al.,  
10 2011; Rimkus, 1999). Here, we examine the dynamics of antibiotic mass  
11 fluxes in a wastewater treatment plant (WTP) influent.

12     Antibiotics have been measured in WTP influents worldwide (Hirsch  
13 et al., 1999; Kümmerer, 2009; Tamtam et al., 2008). They are suspected to  
14 present an environmental risk, e.g., fostering bacterial resistance (Czekalski  
15 et al., 2010; Servais and Passerat, 2009). For this reason, antibiotics in ur-  
16 ban systems and discharges have been subject to increased investigation in  
17 the last decade (Neu, 1992; Pruden et al., 2006; Verlicchi et al., 2012). Sev-  
18 eral studies have reported evidence of seasonal fluctuations in concentra-  
19 tions of various antibiotics in WTPs and in natural waters (Conley et al.,  
20 2008; Göbel et al., 2005; Lissemore et al., 2006; Santos et al., 2009). The  
21 changing wastewater characteristics and imposition of stricter wastewater  
22 discharge regulations have led to a greater emphasis on WTP inlet fluxes  
23 (Tchobanoglous et al., 2003). Process modeling of activated sludge dynam-  
24 ics is becoming increasingly important, as is the characterization of the dy-  
25 namics of WTP influent (Hulsbeek et al., 2002). The temporal dynamics of

26 ammonia and dissolved organic carbon for instance have already been well  
27 described (Tchobanoglous et al., 2003). However, there are few studies that  
28 assess the time variability of antibiotics. In Table 1, we summarize existing  
29 studies reporting antibiotic dynamics at WTP inlets.

30 In a paper aiming to assess the removal of pharmaceuticals and fragrances  
31 in a WTP, Joss et al. (2005) observed some fluctuations in antibiotic con-  
32 centrations at the WTP inlet, but the sampling methodology (8-h composite  
33 samples, Table 1) was insufficient to determine details of the influent dy-  
34 namics. In another study, Plósz et al. (2010) measured the diurnal variation  
35 of hormones and antibiotics in activated sludge wastewater, as part of their  
36 overall goal of linking posology and observed concentrations. However, sim-  
37 ilarly to Joss et al. (2005), Plósz et al. (2010) used 8-h composite samples  
38 taken after the primary clarifier to assess the temporal variability (Table 1).  
39 That is, in both studies, the authors collected three 8-h composite samples  
40 per day. Plósz et al. (2010) had a total of nine samples (3 sampling days)  
41 whereas Joss et al. (2005) sampled for 1 d giving a total of three samples. In  
42 both cases, the sampling frequency (3 per day) and the total number of sam-  
43 ples are insufficient to identify diurnal variability. Gerrity et al. (2011) and  
44 Teerlink et al. (2012) identified a diurnal pattern consisting of two mass flux  
45 peaks in WTP influent, one in the morning followed by a lower peak during  
46 the night (Table 1). Salgado et al. (2011) investigated day-to-day variation of  
47 chemicals and measured a load of pharmaceuticals that surprisingly varied  
48 by an order of magnitude from one day to the next. These authors con-  
49 cluded that the dynamics of pharmaceuticals needed further investigation to  
50 understand the variability in more detail.

51 **Table 1 near here**

52 The studies listed in Table 1 generally observed a peak in morning con-  
53 centrations. This observation is not surprising as antibiotics and other sub-  
54 stances ingested during the evening accumulate in urine during sleep (Noh  
55 et al., 2011; Pandi-Permural and Cardinali, 2007). However, all the cited  
56 studies were not designed to examine the temporal variability of antibiotics  
57 in WTP influent in a systematic way (i.e., over different time scales). In this  
58 work, our objective was to uncover time scales in mass fluxes of represen-  
59 tative antibiotics in WTP influent. To this end, we performed a yearlong  
60 on-site experimental campaign with sufficient sampling to capture hourly to  
61 seasonal fluctuations. Besides its scientific interest, the study has the poten-  
62 tial to help in the optimization of micropollutant removal in WTPs. More  
63 particularly, the results will underpin the design of future sampling cam-  
64 paigns focusing on antibiotics, as well as the interpretation of data sets, e.g.,  
65 short-term sampling campaigns.

## 66 **2. Materials and Methods**

### 67 *2.1. Sample collection*

68 A yearlong study was conducted at the Vidy WTP, in Lausanne, Switzer-  
69 land. The WTP collects wastewater for approximately 220,000 inhabitants,  
70 and several hospitals. The latter contribute about  $\sim 1\%$  of the total wastew-  
71 ater volume of reaching the WTP. The automatic sampler was installed at  
72 the entrance of the WTP, upstream of the primary settling tank, in order to  
73 capture the dynamics of the mass flux of incoming antibiotics. Starting in

74 March 2011, 12 (i.e., once each month) weeklong sampling campaigns were  
75 conducted. These campaigns consisted of seven daily-composite samples,  
76 collected for the seven consecutive days (84 samples collected in total). A  
77 daily-composite sample was obtained by mixing (flow proportionally) 24 indi-  
78 vidual (hourly) 200-ml samples collected with an automatic sampling device  
79 (6712FR Teledyne ISCO). Collected samples were stored on-site in plastic  
80 bottles inside a refrigerated sampling device at a temperature below 4°C be-  
81 fore collection for laboratory analysis. Samples were analyzed or frozen at  
82 -20°C within 24 h after collection.

83 In addition to the yearlong campaign, four 24-h campaigns were con-  
84 ducted to evaluate the dynamics at the hourly scale. The four campaigns  
85 were conducted during four different months: December, May, September  
86 and November. Each 24-h campaign provided 24 samples (96 in total), one  
87 per hour, starting at 07:00. Each hourly sample was the aggregation of four  
88 200-ml wastewater samples collected every 15 min.

89 The different field campaigns are summarized in Table 2. All samples were  
90 collected at the same location using the same sampler. Water samples were  
91 analyzed using online SPE UPLC-MS/MS, and concentrations were obtained  
92 for six antibiotics — trimethoprim, norfloxacin, ciprofloxacin, ofloxacin, clin-  
93 damycin, metronidazole — for a total of 1080 analyses. These six antibiotics  
94 were classified as priority substances based on physico-chemical characteris-  
95 tics in a previous study (Coutu et al., 2012) and because they were measured  
96 in Vidy Bay, where the WTP discharges its effluent (Bonvin et al., 2011).

97 The sampling program allows comparison of mass flux fluctuations month-  
98 to-month, day-to-day and hour-to-hour, thereby providing information on the

99 time scales that control antibiotic fluxes in wastewater.

100 **Table 2 near here**

101 *2.2. Flow measurements*

102 An ultrasonic sensor was installed next to the automatic sampler to record  
103 water levels at 5-min intervals. The water level was later converted into flow  
104 by use of the appropriate calibration<sup>1</sup>. All water flow data can be found in  
105 the Supplementary data.

106 *2.3. Sample treatment and chemical analysis*

107 As a first step samples were acidified to pH 2-2.5 with hydrochloric acid  
108 (concentrated 25%). Then, samples were filtered, first, through 2.7- $\mu$ m glass  
109 fiber pre-filters (type GF/D, Whatman). The pre-filtered samples were fil-  
110 tered again through 0.45- $\mu$ m membrane filters (type ME 25, mixed cellulose  
111 ester, Whatman). Samples were stored at -20°C until analysis.

112 The analytical method involved online solid-phase extraction (SPE) and  
113 Ultra Performance Liquid Chromatography coupled with tandem mass spec-  
114 trometer (Xevo UPLC-MS/MS, Waters). Frozen samples were thawed to  
115 room temperature prior to analysis. Then, 8 mL of each sample was fil-  
116 tered through 0.2- $\mu$ m syringe filters (type GMF, BGB-analytik) directly into  
117 the injection vials. A mixture of deuterated antibiotic surrogates was added  
118 with a gas-tight syringe into all injection vials (samples as well as standards).  
119 Targeted compounds were first extracted in the online SPE system, which

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<sup>1</sup><http://www.isco.com>, site last accessed in June 2012

120 consisted of a 2.1×20 mm SPE column (type Oasis HLB 25  $\mu\text{m}$ , Waters).  
121 Extracted compounds were separated in a 2.1 × 50 mm chromatographic  
122 analytical column (type Acquity UPLC BEH C18 1.7  $\mu\text{m}$ , Waters) with an  
123 organic mobile phase in gradient mode. All the targeted compounds were  
124 identified and quantified using tandem mass spectrometry according to their  
125 masses of precursor and product ions as well as their mass-to-charge ratios.  
126 More information on mobile phases and HPLC gradient used can be found  
127 in the supplementary materials.

128 The analytical limit of quantification (LOQ) was defined as the concen-  
129 tration of the lowest standard with a signal-to-noise ratio greater than 10  
130 (Bonvin et al., 2011). The antibiotic concentration in the samples was calcu-  
131 lated based on calibration curves using seven calibration points closest to the  
132 sample concentration. Correlation coefficients for the calibration curves were  
133 typically 0.99 at least. In the calculation of sample concentration, recovery  
134 rates of deuterated surrogates and exact sample mass weighed during sample  
135 preparation were taken into account for each associated antibiotic compound.  
136 Details of the precision of the method are summarized in Table 3.

#### 137 *2.4. Data analysis*

138 We discuss mass fluxes of antibiotics rather than concentrations, since  
139 dilution occurs during rainy periods. Mass fluxes, or load, to the WTP  
140 is useful when different sites are compared (Johnson, 2010). Hereafter, we  
141 use the term “measured mass flux” to indicate the mass flux obtained by  
142 multiplying, for each sample, the measured flow and measured concentration.

143 Statistical techniques were used to assess potential temporal character-  
144 istics in mass flux fluctuations and the variability at different time scales

145 (month, day, hour: m, d, h) in the mass flux of antibiotics at the WTP inlet.  
146 Calculations were carried out using Matlab<sup>2</sup>. Standard statistical tests were  
147 performed (mean, median and different percentiles of the measured mass  
148 fluxes). In addition, the periodogram of the 96 hourly values was extracted  
149 to identify possible periodicity in the hourly dynamics. This was done by  
150 windowing the data using the technique of Welch (1967), after which the  
151 periodogram was computed using the fast Fourier transform.

### 152 **3. Results and Discussion**

#### 153 *3.1. Seasonality of mass flux*

154 Mass fluxes measured monthly at the WTP inlet are compared. Seven  
155 consecutive daily samples were aggregated flow proportionally to form a  
156 monthly sample (Table 2). The median mass flux obtained is presented for  
157 each month and all substances in Figure 1. Minimum monthly mass fluxes are  
158 generally found during the period running from June to September. Mass  
159 fluxes measured in December and January have systematically higher val-  
160 ues except for ciprofloxacin. High values are also observed in March for  
161 ciprofloxacin, ofloxacin and clindamycin. Note that an error in the experi-  
162 mental procedure meant that unfortunately we could not retrieve mass fluxes  
163 for the month of February.

164 In comparing the different time series of mass fluxes, data for a given time  
165 scale were normalized by the appropriate mean calculated for the considered  
166 data. Thus, hourly, weekly and monthly data were normalized by daily,

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<sup>2</sup>Matlab 2009b, The Mathworks, Natick, Massachusetts, USA

167 monthly and annual means, respectively. This allows comparison of varia-  
168 tions in data sets with different means. For example, the monthly mass flux  
169 normalized by the annual mean is helpful to evaluate the seasonality of the  
170 different antibiotics, as well as the range of the month-to-month fluctuations.  
171 This relative fluctuation is presented in Figure 3a for the six antibiotics. The  
172 normalized monthly mass fluxes range between a quarter and double of the  
173 mean annual mass flux measured at the WTP inlet. Obvious seasonality  
174 is observed in the mass fluxes of ciprofloxacin, ofloxacin and clindamycin,  
175 with winter values being up to eight times greater than corresponding sum-  
176 mer values. Trimethoprim, on the other hand, shows less variability in the  
177 monthly mass flux over the year, although a higher mass flux is still observed  
178 in January and December. The behavior of norfloxacin and metronidazole  
179 is somewhat peculiar; values of monthly mass fluxes are located broadly be-  
180 tween a quarter and half of the mean, with a sudden rise in December and  
181 January. For all substances we observe that the highest mass flux is measured  
182 during the (winter) months of December and January.

183 The boxplot widths appear to be significant for some substances in Fig-  
184 ure 1 (e.g., norfloxacin in January). This indicates that successive daily mass  
185 fluxes within a week were not constant, which should be the case for antibi-  
186 otics (see §.3.3). This variability reflects the random nature of the inputs  
187 into the sewer system, as well as uncertainty in the measurements, which in  
188 the case of UPLC/MS-MS can be reasonably considered to range up to 20%  
189 (Bonvin et al., 2011).

190 **Figure 1 near here**

191 The clear seasonality in the observed mass flux of ciprofloxacin (Figure 1)  
192 can be explained by considering its therapeutic use. Ciprofloxacin is mainly  
193 prescribed to treat seasonal diseases like airway infections (i.e., bronchitis and  
194 pneumonia) and infections affecting the throat, nose and ears (pharyngitis,  
195 sinusitis, earache). The mass flux of norfloxacin is also affected by seasonality.  
196 It is also used to treat similar seasonal pathologies, albeit to a lesser extent  
197 than ciprofloxacin. In contrast, trimethoprim displays reduced seasonality  
198 in mass fluxes, corresponding to its use for non-seasonal diseases, typically  
199 urinary-genital tract infections. However, the connection between seasonality  
200 in prescriptions to that in mass flux is not systematic. Indeed, ofloxacin  
201 displays seasonality in mass flux dynamics, yet it is used to treat non-seasonal  
202 pathologies like skin or urinary-genital tract infections. It is thus difficult  
203 to explain seasonality (or lack thereof) in mass fluxes of specific compounds.  
204 Possibly, seasonal temperature changes are sufficient to influence degradation  
205 of some compounds while they are transported in the sewer system.

### 206 *3.2. Periodicity in hourly mass flux*

207 The four 24-h campaigns described in Table 2 were used to evaluate peri-  
208 odicity in the mass flux of antibiotics reaching the WTP during dry weather.  
209 For this, the measured hourly mass flux was first normalized to the mean for  
210 that day (Figure 3c). After normalization, data from the four 24-h campaigns  
211 were combined sequentially into a series of 96 experimental values. The pe-  
212 riodogram for ciprofloxacin is shown in Figure 2. Periodograms for the five  
213 other antibiotics considered in this study can be found in the Supplementary  
214 data.

215 **Figure 2 near here**

216 For all substances considered, two noticeable peaks appear in the com-  
217 puted periodogram, which indicates that the mass flux of antibiotics has a  
218 12-h period. According to Figure 3c, the first peak occurs between 07:00 and  
219 09:00. The magnitude of this morning peak is up to double that of the aver-  
220 age daily flux reaching the WTP. It corresponds to the peak in toilet flushes  
221 observed by Friedler et al. (1996) and Rauch et al. (2003). The second peak  
222 occurs 12 h later, between 17:00 and 21:00. This peak is less pronounced  
223 than the morning peak, and occurs over a broader period. Its maximum is  
224 around 1.2 times the daily mean, except for metronidazole.

225 Consistent with previous observations, the fluctuations in the mass flux  
226 of antibiotics are greater than those of the flow rate (Figure 3c). Plósz et al.  
227 (2010) explained that the high peak in pharmaceutical load at the WTP in-  
228 let is due to posology. Antibiotics are typically administrated every 8 or 12  
229 h, with one ingestion occurring in late evening. During sleep, the antibiotic  
230 accumulates in urine and is released in the first toilet flush of the morning.  
231 Concentrations are thus relatively high, leading to the observed difference in  
232 fluctuations. A second peak in the mass flux, lower than the first, occurs ap-  
233 proximately 12 h later (Figure 2 and 3c). This second peak was not observed  
234 by Plósz et al. (2010) as they used 8-h composite samples. The reduction in  
235 the magnitude of this peak results again from posology. Indeed, the posol-  
236 ogy of antibiotics is calculated so that they are metabolized at a constant  
237 rate. Due to more frequent toilet use during daytime, the concentration in  
238 the urine is, ideally, constant. Thus, the second peak observed is due to the  
239 corresponding increase in wastewater volume. For this reason the increase in

240 mass flux of antibiotics and flow rate are in the same range (Figure 3c).

### 241 3.3. Dominant time scale

242 Mean normalized mass fluxes were used to compare the temporal dy-  
243 namics of antibiotics at different time scales. The results are summarized in  
244 Figure 3. We observe that whereas monthly and hourly mean mass fluxes  
245 fluctuate between 0.25 and 2 times the corresponding mean (the annual mean  
246 and daily mean, respectively), very little fluctuation is observed at the daily  
247 scale (Figure 3b). All daily measured mass fluxes remain located between  
248  $\pm 20\%$  of the weekly average, the only exception being metronidazole. A  
249 day-to-day variability in the usage of this last substance could be responsi-  
250 ble for the observed variability at the WTP entrance (between 2.2 and 0.3  
251 times the mean daily load flux, see Figure 3b). Indeed, some pathologies are  
252 treated by a single high dose of this compound (e.g., bacterial vaginosis, 2 g  
253 of metronidazole) (CDC, 2002). Depending on the number of people treated,  
254 these “single pulses” may explain the observed variability at this time scale.  
255 More medical data on the occurrence of these kind of diseases would help to  
256 characterize this behavior.

257 **Figure 3 near here**

258 **Figure 4 near here**

259 The dominance of variability in antibiotic mass fluxes at the monthly and  
260 hourly time scales over that at the daily time scale is illustrated in Figure 4.  
261 The variability of the daily mass flux (during the week) is systematically

262 lower than that of the monthly mass flux considered over the year, or hourly  
263 mass flux over a day. Depending on the month or the hour of the sampling,  
264 the range of mass fluxes varies by up to an order of magnitude. This is  
265 due to two factors. First, antibiotic consumption is seasonal, thus so are  
266 the corresponding mass fluxes at the WTP entrance (Vernaz et al., 2008).  
267 Second, on the hourly time scale (during a given day), mass flux variability  
268 can be (partially) explained by toilet use frequency and the posology of the  
269 different antibiotics. These variations highlight the importance of the design  
270 of sampling campaigns for antibiotics in the environment Ort et al. (2010).

#### 271 *3.4. Comparison with existing studies*

272 Comparisons of these results with existing studies is not straightforward  
273 as different analytical methods have been employed. In addition, there is  
274 variability in use of antibiotics and water between regions (Filippini et al.,  
275 2006).

276 Santos et al. (2009) conducted a 1-y field campaign at a WTP entrance  
277 to assess the dynamics of some micropollutants (not antibiotics). No clear  
278 pattern in the dynamics of the substances considered could be identified  
279 except for ibuprofen, where the authors suspected some seasonality. How-  
280 ever, ibuprofen is an anti-inflammatory drug and seasonality in its use seems  
281 unlikely, especially when contrasted with antibiotics used to treat seasonal  
282 diseases such as airway infections (i.e., bronchitis and pneumonia) and in-  
283 fections affecting the throat, nose and ears (pharyngitis, sinusitis, earache).  
284 For the Lausanne WTP, ciprofloxacin shows the clearest seasonality, with the  
285 mass flux in winter being up to 4 times greater than that measured during  
286 summer.

287 Göbel et al. (2005) assessed the seasonality of different antibiotics by com-  
288 paring three weekly campaigns conducted in March, September, November  
289 and February (see Table 1) in Eastern Switzerland. Loads observed at the  
290 WTP inlet were generally higher in February and March than in Septem-  
291 ber and November. This was explained by higher sales during these periods  
292 (sales data not provided) and, as mentioned above, by the seasonal use of  
293 these substances determined by the treated pathologies. Unfortunately, the  
294 number of samples collected was insufficient to determine a clear pattern in  
295 seasonal fluctuations of loads.

296 Salgado et al. (2011) assessed diurnal variability of several pharmaceuti-  
297 cals (and other chemicals) in wastewater influent. They sampled wastewater  
298 at a WTP inlet during two consecutive days (Monday-Tuesday), two weeks  
299 in a row. Ciprofloxacin was measured on the two Tuesdays. A factor of  
300 two in mass load was observed, which was difficult to explain given that the  
301 sampling methodology and weather conditions were consistent for the two  
302 sampling days. By comparison, our results have clearer underpinnings, due  
303 to the nested temporal sampling employed. For antibiotics, where the patient  
304 follows a prescribed posology, the consumption over the whole WTP basin  
305 would vary little from one day to the next. Thus, the daily mass flux should  
306 remain within the same order of magnitude. Consistent with this reasoning  
307 are the findings of Gerrity et al. (2011) and Huerta-Fontela et al. (2008), who  
308 found that compounds not subject to any posology (e.g., illicit products) do  
309 not follow this behavior.

310 The diurnal variability of pharmaceuticals was recently the subject of a  
311 study conducted at a WTP inlet in Norway. Plósz et al. (2010) observed a

312 mass load up to three times greater in the morning than during the night  
313 for ciprofloxacin and trimethoprim. A similar pattern was observed in other  
314 studies assessing the temporal dynamics of pharmaceuticals (Göbel et al.,  
315 2005; Joss et al., 2005). A second peak observed by Teerlink et al. (2012)  
316 and Gerrity et al. (2011) in the late afternoon was simply explained by a  
317 corresponding rise in flow rate at WTP inlet. However, it would be beneficial  
318 to combine knowledge of posology, pharmacokinetics and toilet use frequency  
319 to obtain more insights on the hourly fluctuations of these substances at the  
320 WTP inlet. A modeling study in that direction is currently underway in our  
321 group.

#### 322 **4. Conclusion**

323 We have assessed the dominant time scale in the mass flux variability of  
324 several antibiotics in sewage water reaching the Vidy WTP. Mass fluxes were  
325 observed to be higher in winter than in summer. This is particularly the  
326 case for ciprofloxacin, with the winter mass flux being up to 3-4 times that  
327 in summer. These data are consistent with the hypothesis that the measure-  
328 ments reflect the seasonal consumption of antibiotics, driven by associated  
329 seasonality in pathologies. However, further investigation is necessary to es-  
330 tablish a correlation between consumption and measured load to confirm this  
331 hypothesis.

332 Little fluctuation was observed on a day-to-day basis. This indicates that,  
333 at the scale of the WTP catchment area, daily consumption of antibiotics  
334 varies only slowly. Metronidazole was found to behave differently, however,  
335 for reasons that are as yet unknown.

336 Diurnal fluctuations in the mass flux of antibiotics show a similar trend  
337 for the six compounds investigated. Similarly to previous studies, a morning  
338 peak in mass flux was observed, which was not surprising as antibiotics accu-  
339 mulate overnight in the human body before being excreted in the morning.  
340 Another peak was generally observed 12 h later. Even though it is plausible  
341 that toilet use could explain this second peak, more knowledge on the phar-  
342 macokinetics of these antibiotics seems necessary to characterize more fully  
343 this mass flux increase.

344 Several conclusions on sampling design of antibiotics in wastewater can be  
345 drawn out of the results of this study. First, the day of the week selected for  
346 sampling has no influence on the load measured. Second, the seasonal con-  
347 sumption of the substance should be investigated when possible, as sizeable  
348 fluctuations of mass flux of antibiotics can be observed at the WTP inlet.  
349 Third, at least an hourly sampling frequency and analysis are needed when  
350 the objective of the study is to assess the intra-day dynamics of antibiotics,  
351 in order to capture the morning and late afternoon mass flux peaks.

352 More generally, posology, pharmacokinetics and toilet flush frequency  
353 control the mass flux of antibiotics in sewage water. These parameters should  
354 be accounted for when modeling the mass flux or concentrations in WTPs.  
355 The dataset presented here will aid such modeling efforts in the future.

## 356 **Supplementary data**

357 Supporting information is provided in the three files:

- 358 • A1 (A1\_flow.xlsx): A spreadsheet (Excel) file containing all flow values  
359 measured during the different experimental campaigns;

- 360 • A2 (A2\_concentrations.xlsx): A spreadsheet (Excel) file containing all  
361 concentrations measured during the different experimental campaigns  
362 for each substance;
- 363 • A3 (A3\_periodo\_all): A PDF file with six figures representing the peri-  
364 odograms computed for the six antibiotics (cf. Figure 2);
- 365 • A4 (A4\_GradientMode.docx): A word file with additional technical  
366 notes on the experimental procedure.

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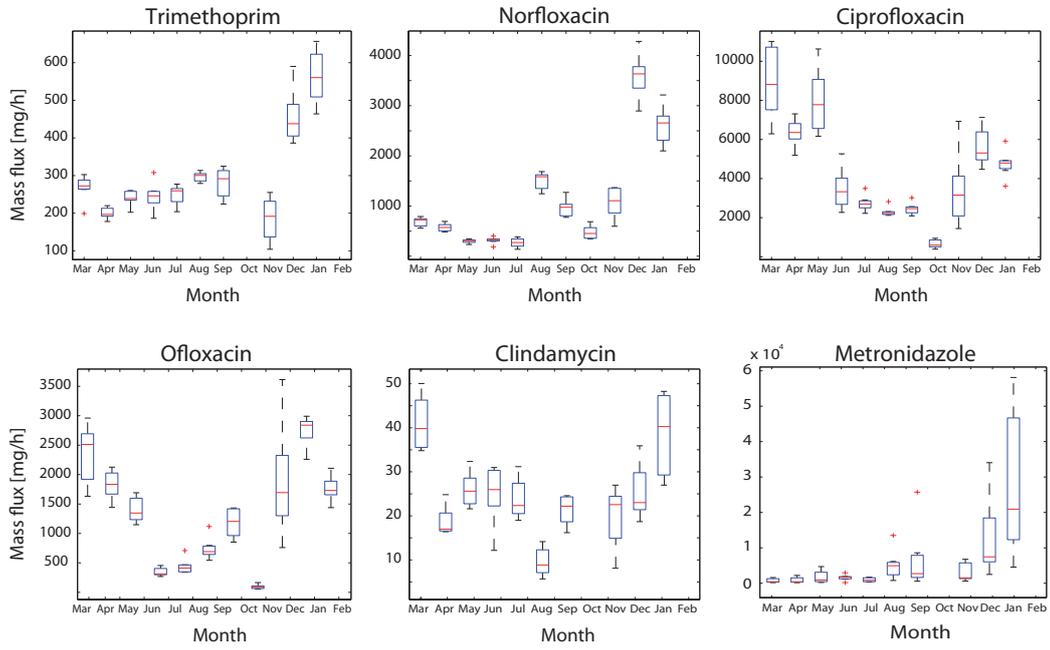


Figure 1: Boxplots of mass fluxes of the six antibiotics investigated (one box per month). Each box was obtained from seven daily samples. Upper and lower box limits correspond to the 75<sup>th</sup> and 25<sup>th</sup> percentiles, respectively. Upper and lower whiskers correspond to the last datum being respectively still within 1.5IQR of the higher and lower quartile. IQR is the interquartile range, i.e., the difference between the upper and lower quartiles.

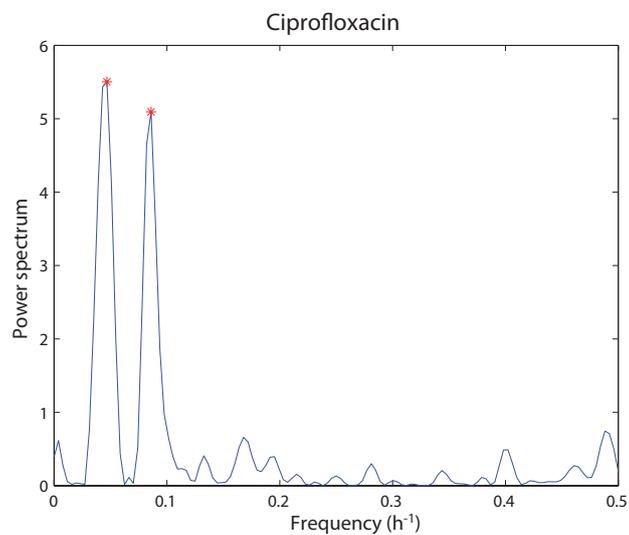


Figure 2: Periodogram of the measured ciprofloxacin hourly mass flux. Peaks in the power spectrum observed at  $0.042 \text{ h}^{-1}$  and  $0.083 \text{ h}^{-1}$ , which correspond to periods of 24 h and 12 h, respectively. The same periods were observed for the other substances (Supplementary data).

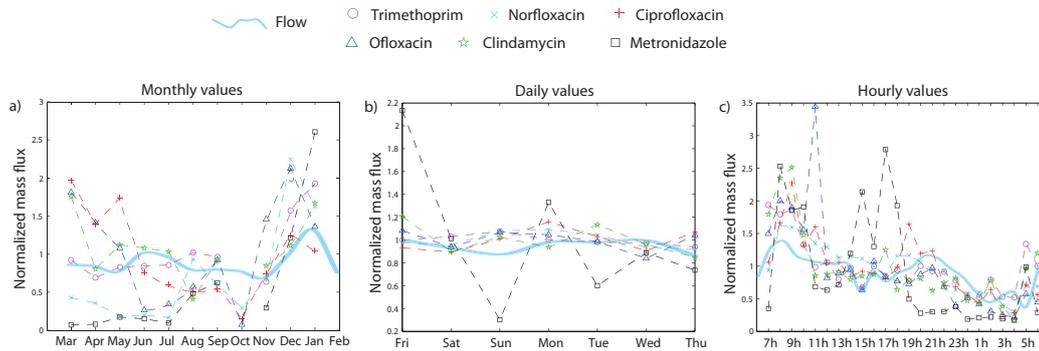


Figure 3: Fluctuations of mass flux at different time scales: month a), day b), and hour c). In a), each symbol corresponds to the average mass flux of the seven daily samples collected in the corresponding month. In b), each symbol corresponds to the average value of the twelve times (one per month) the corresponding day was sampled. In c), each point corresponds to the average of the four times (four 24-h campaigns) the corresponding hour was sampled. Mass flux is normalized to mean mass flux over the considered time scale. For example, in a) the monthly mass flux was normalized by the mean of the monthly mass fluxes measured over the year. Clear variability is observed at the monthly and hourly time scales for all substances. No variability is observed at the daily time scale, except for metronidazole. The corresponding flow — normalized to the mean flow over the considered time scale — is shown on each graph by the blue line. The monthly mean flow includes rainwater peaks in addition to baseflow. Daily and hourly mean flows were selected to include only dry periods, i.e., their fluctuation is not affected by any rain input.

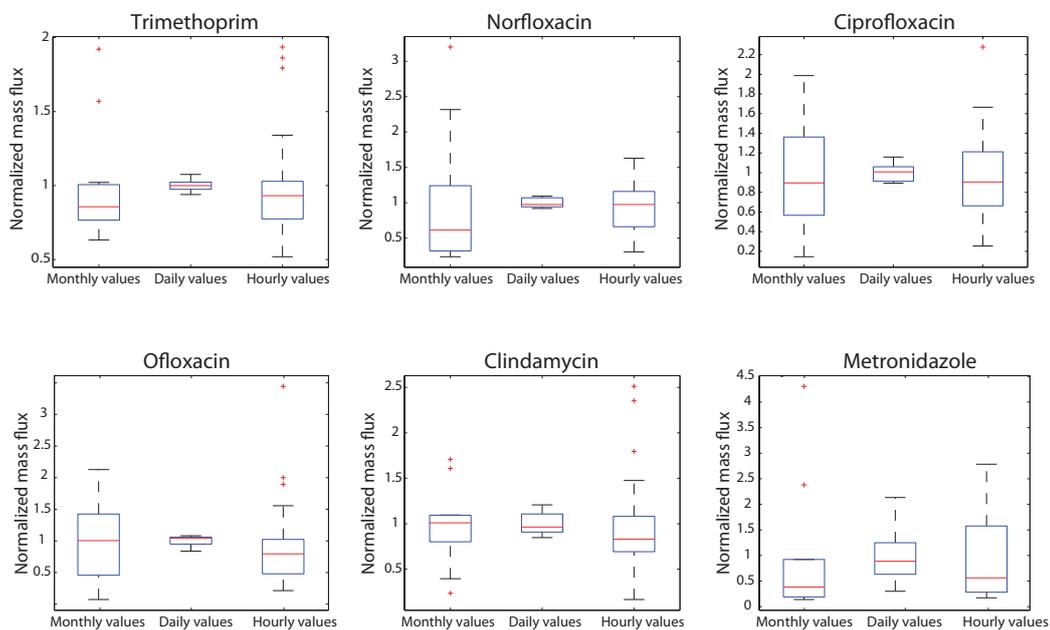


Figure 4: Boxplots illustrating the dispersion of the measured mass fluxes for the three different time scales (m, d, h). Similar to Figure 3, the mass flux is normalized by the mean mass flux over the considered time scale. For each substance, the box on the left is obtained from the 12 mean monthly values (the 12 values in Figure 3a), the box in the middle from the seven mean daily values (seven values in Figure 3b), and the box in the right from the 24 mean hourly values (the 24 values in Figure 3c). Boxes were computed following the same rule described in Figure 1.

Table 1: Summary of studies assessing the temporal dynamics of antibiotics in WTP influent.

<b>Study</b>	<b>Antibiotic</b>	<b>Sampling regime at the WTP inlet</b>	<b>Sampler position</b>
Joss et al. (2005)	Roxithromycin, sulfamethoxazole	8-h composite samples of primary effluent taken during one day	After primary clarifier
Göbel et al. (2005)	Sulfadiazine, sulfathiazole, sulfapyridine, sulfamethoxazole, trimethoprim, erythromycin, clarithromycin	Flow-proportional 8-h composite sample during one day plus three weekly campaigns to assess seasonality (three flow-proportional composite samples, one combining 24-h integrated samples from Saturday and Sunday and two combining two or three weekdays)	WTP inlet, before clarifier
Plósz et al. (2010)	Ciprofloxacin, trimethoprim, sulfamethoxazole, tetracycline	Three-day measurement campaign. 8-h flow-proportional composite samples. Sampling frequency of 15 min	After primary clarifier
Ort et al. (2010)	Chloramphenicol, trimethoprim, roxithromycin, lincomycin, erythromycin, cephalexin, dapson, sulphadiazine	Tested several sampling strategies and recommended a 5-min sampling frequency to characterize intra-day fluctuations in WTP influent	WTP inlet, before clarifier
Gerrity et al. (2011)	Trimethoprim, sulfamethoxazole	Two 12-h campaigns with a sampling frequency of 30 min	After primary clarifier
Salgado et al. (2011)	Ampicillin	Two 48-h campaigns with a sampling frequency of 2 h	WTP inlet, before clarifier
Teerlink et al. (2012)	Trimethoprim, sulfamethoxazole	One 26-h campaign with a sampling frequency of 1 h	WTP inlet, before clarifier

Table 2: Description of the methodology used to obtain the monthly, daily and hourly samples in the field campaign. Samples were all collected at the same location at the Vidy WTP entrance.

Type of sample	Sample aggregation	Number of samples	Frequency
Monthly samples	Flow proportional average of seven consecutive daily samples	12	One per month
Daily samples	200 ml samples collected every hour during a day. The 24 samples were mixed flow proportionally to get the daily average	84	7 d (1 week) per month
Hourly samples	200 ml samples collected every 15 min during an hour. Each hour, the four 200 ml samples were aggregated to form an hourly sample	96	Four 24 h campaigns (May, September, November, December)

Table 3: Precision of the laboratory analysis.

	Analyte	RSD (%) <sup>a</sup>		LOD <sup>c</sup> (ng/L)	LOQ <sup>d</sup> (ng/L)	Recovery (%) <sup>e</sup>
		Within one run	(R <sup>2</sup> ) <sup>b</sup>			
1	Trimethoprim	2	0.9993	1.3	4	88±1
2	Norfloxacin	8	0.9827	1.3	4	118±43
3	Ciprofloxacin	1	0.9963	1.2	3.5	105±7
4	Ofloxacin	7	0.9959	1.2	3.5	90±11
5	Clindamycin	1	0.9961	1.1	3.3	86±1
6	Metronidazole	4	0.9972	1	3	116±22

<sup>a</sup>Relative Standard Deviation: concentrations detected in the samples during one continuous sequence of analysis. Three replicates were carried out

<sup>b</sup>Correlation coefficient with seven concentration levels. Norfloxacin, ofloxacin: 0.2 - 350 ng/L; trimethoprim, clindamycin: 0.3 - 630 ng/L and ciprofloxacin, metronidazole: 0.4 - 700 ng/L

<sup>c</sup>Limit of detection calculated as a signal to noise (S/N) ratio of 3/1

<sup>d</sup>Limit of quantification calculated as a signal to noise (S/N) ratio of 10/1

<sup>e</sup>Quality control samples (three replicates) prepared in three injection vials for each analyte with known concentrations of 3, 30 and 150 ng/L. Three replicates were carried out